

Table EV1 related to Figure 1.

Statistics of transcriptome and proteome changes during *N. furzeri* brain aging.

Protein groups			
	12 vs. 5 stable	12 vs. 5 up	12 vs. 5 down
39 vs. 12 stable	3462	493	837
39 vs. 12 down	690	111	252
39 vs. 12 up	978	127	250

Out of 7200 detected proteins (overlap between the two conditions).

Transcripts			
	12 vs. 5 stable	12 vs. 5 up	12 vs. 5 down
39 vs. 12 stable	18557	605	244
39 vs. 12 down	786	63	25
39 vs. 12 up	885	144	92

Out of 21813 transcripts (overlap between the two conditions).

Legend

Stable = not significant
Down = adj. $p < 0.05$ & $\log_2 FC < 0$
Up = adj. $p < 0.05$ & $\log_2 FC > 0$